**In the Figures:**

Please delete the disclosure on page 46 and replace it with the following:

DEFINITION Human Cockayne syndrome complementation group A CSA protein (CSA) mRNA, complete cds. **(SEQ ID NO: 11)**
ACCESSION U28413

BASE COUNT 596 a 368 c 413 g 634 t
ORIGIN

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1       CGACGTCCAG TGCTCCAGCC GGTGTGAGGA CACGATATGC TGGGGTTTTT GTCCGCACGC
61      CAAACGGGTT TGGAGGACCC TCTTCGCCTT CGGAGAGCAG AGTCAACACG GAGAGTTTTG
121     GGACTGGAAT TAAATAAAGA CAGAGATGTT GAAAGAATCC ACGGCGGTGG AATTAACACC
181     CTTGACATTG AACCTGTTGA AGGGAGATAC ATGTTATCAG GTGGTTCAGA TGGTGTGATT
241     GTACTTTATG ACCTTGAGAA CTCCAGCAGA CAATCTTATT ACACATGTAA AGCAGTGTGT
301     TCCATTGGCA GAGATCATCC TGATGTTCAC AGATACAGTG TGGAGACTGT ACAGTGGTAT
361     CCTCATGACA CTGGCATGTT CACATCAAGC TCATTTGATA AAACTCTGAA AGTATGGGAT
421     ACAAATACAT TACAAACTGC AGATGTATTT AATTTTGAGG AAACAGTTTA TAGTCATCAT
481     ATGTCTCCAG TCTCCACCAA GCACTGTTTG GTAGCAGTTG GTACTAGAGG ACCCAAAGTA
541     CAACTTTGTG ACTTGAAGTC TGGATCCTGT TCTCACATTC TACAGGGTCA CAGACAAGAA
601     ATATTAGCAG TTTCCTGGTC TCCACGTTAT GACTATATCT TGGCAACAGC AAGTGCTGAC
661     AGTAGAGTAA AATTATGGGA TGTGAGAAGA GCATCAGGAT GTTTGATTAC TCTTGATCAA
721     CATAATGGGA AAAAGTCACA AGCTGTTGAA TCAGCAAACA CTGCTCATAA TGGGAAAGTT
781     AATGGCTTAT GTTTTACAAG TGATGGACTT CACCTCCTCA CTGTTGGTAC AGATAATCGA
841     ATGAGGCTCT GGAATAGTTC CAATGGAGAA AACACACTTG TGAACTATGG AAAAGTTTGT
901     AATAACAGTA AAAAAGGATT GAAATTCACT GTCTCCTGTG GCTGCAGTTC AGAATTTGTT
961     TTTGTACCAT ATGGTAGCAC CATTGCTGTT TATACAGTTT ACTCAGGAGA ACAGATAACT
1021    ATGCTTAAGG GACATTATAA AACTGTTGAC TGCTGTGTAT TTCAGTCAAA TTTCCAGGAA
1081    CTTTATAGTG GTAGCAGAGA CTGCAACATT CTGGCTTGGG TTCCATCCTT ATATGAACCA
1141    GTTCCTGATG ATGATGAGAC TACAACAAAA TCACAATTAA ATCCGGCCTT TGAAGATGCC
1201    TGGAGCAGCA GTGATGAAGA AGGATGAATA TCATCTTTAG TACCTTTTTG TCTCTGCTGA
1261    AACTTTTTAA ATGAGACTGT GTTTTTTTCA ACTGTATGGT CTATTCCTGA CAGCTAAATT
1321    AGCCCTAAAT GCGGGTAATA TTTTTCCTCA TGTTTTAAAA TGAGGTTAAT ATTTGCATAA
1381    AATCCTAAAA CAGACTTCTG TATAGTTTAT TTAGTCAAAA TGTGTTCCTT GATCCCAGAT
1441    GTTGTGGCCT GGGAAAGCCC TCATTGCTAC AGTACAAGTA ACACAAGTCG TTGTACCTCA
1501    GTTGTGACCT TCAGCAGATT TTATGAACTA TAAGATGCAG TCTCAGAGGA TCAGCAAGTG
1561    GAGGCCATCA GTATTGACTT TCTCTTACTT GCTGTACTAT CAGCCTGCTC GTTTCCACCT
1621    TTAAGAATGA TTTTGCCAAG AATGATTATA TCAAAAATAG TAGTTGAAAT GGTAACATCA
1681    AAATTATTTT ATTCTTTCTT CTTCATGTAT TCACATTTTT CAGTGGTTTC ATTTAATTAA
1741    CCATGCTTTA TGTTAAACAT TTTGGGGCTC AATGTCTCCT ACTATCCAAA ATGTGCATCA
1801    CAGGAGGCTC TTAACTTTGT GAAAATCCCA TGTTTGCTTT ATTTTATTTT AATGTCAGAA
1861    GGCAGTTTGC GCTAATGCTT GAACTCTTTT TCTGTGAAAC TCATTAAGGT ATGACCAAAT
1921    CCTGCCTCAT TAATTCAAGC AGAAAATATC CTGGCAGGGA ATCTGGCTTA AACATGAAAT
1981    GCTGTAATAA AATTTCTATG TTATTGTCTC A
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Please delete the disclosure on page 47 and replace it with the following:

DEFINITION Human excision repair protein ERCC6 mRNA, complete cds. **(SEQ ID NO: 12)** (CSB protein)
ACCESSION L04791

BASE COUNT 1433 a 993 c 1220 g 1068 t
ORIGIN

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1       TGGGTTCCAA GGCGGCTGGC GGCGGTAGCG TCTCTGTTTC CTTGTGGGCG CTCGCGCGGC
61      CCTGGGTAGT CTGTAGAGAA TGCCAAATGA GGGAATCCCC CACTCAAGTC AAACTCAGGA
121     GCAAGACTGT TTACAGAGTC AACCTGTCAG TAATAATGAA GAAATGGCAA TCAAGCAAGA
181     AAGTGGTGGT GATGGGGAGG TGGAGGAGTA CCTGTCCTTT CGTTCTGTGG GTGACGGGCT
241     GTCCACCTCT GCTGTGGGGT GCGCATCAGC AGCTCCGAGG AGAGGGCCAG CCCTGCTGCA
301     CATCGACCGA CATCAGATCC AGGCAGTAGA GCCTAGCGCC CAGGCCCTTG AGCTGCAGGG
361     TTTGGGTGTG GACGTCTATG ACCAGGACGT GCTGGAACAG GGAGTGCTTC AGCAGGTGGA
421     CAATGCCATC CATGAGGCCA GCCGTGCCTC CCAGCTCGTT GACGTGGAGA AGGAGTATCG
481     GTCGGTCCTG GATGACCTCA CGTCATGTAC GACATCCCTA AGGCAAATCA ATAAAATTAT
541     TGAACAGCTT AGCCCTCAAG CTGCCACCAG CAGAGACATC AACAGGAAAC TAGATTCTGT
601     AAAACGACAG AAGTATAATA AGGAACAACA GCTAAAAAAG ATCACTGCAA AACAAAAGCA
661     TCTCCAGGCC ATCCTTGGAG GAGCAGAGGT GAAAATTGAA CTAGATCACG CCAGTCTGGA
721     GGAGGATGCA GAGCCGGGGC CATCCAGTCT TGGCAGCATG CTCATGCCTG TCCAGGAGAC
781     TGCCTGGGAA GAGCTCATCC GCACTGGCCA GATGACACCT TTTGGTACCC AGATCCCTCA
841     GAAACAGGAG AAAAAGCCCA GAAAAATCAT GCTTAATGAA GCATCAGGCT TCGAAAAGTA
901     TTTGGCAGAT CAAGCAAAAC TGTCTTTTGA AAGGAAGAAG CAAGGTTGTA ATAAAAGAGC
961     AGCTAGAAAA GCTCCAGCCC CAGTCACGCC TCCAGCCCCA GTGCAAAATA AAAACAAACC
1021    AAACAAGAAA GCCAGAGTTC TGTCCAAAAA AGAGGAGCGT TTGAAAAAGC ACATCAAGAA
1081    ACTCCAGAAG AGGGCTTTGC AGTTCCAGGG GAAAGTGGGA TTGCCAAAGG CAAGGAGACC
1141    TTGGGAGTCA GACATGAGGC CAGAGGCAGA GGGAGACTCT GAGGGTGAAG AGTCTGAGTA
1201    TTTCCCCACA GAGGAGGAGG AAGAGGAGGA AGATGACGAG GTGGAGGGGG CAGAGGCGGA
1261    CCTGTCTGGA GATGGTACTG ACTATGAGCT GAAGCCTCTG CCCAAGGGCG GGAAACGGCA
1321    GAAGAAAGTG CCAGTGCAGG AGATTGATGA TGACTTTTTC CCAAGTTCTG GGGAAGAAGC
1381    TGAAGCTGCT TCTGTAGGAG AAGGAGGAGG AGGAGGTCGG AAAGTGGGAA GATACCGAGA
1441    TGATGGAGAT GAAGATTATT ATAAGCAGCG GTTAAGGAGA TGGAATAAAC TGAGACTGCA
1501    GGACAAAGAG AAACGTCTGA AGCTGGAGGA CGATTCTGAG GAAAGTGATG CTGAATTTGA
1561    CGAAGGTTTT AAAGTGCCAG GTTTTCTGTT CAAAAAGCTT TTTAAGTACC AGCAGACAGG
1621    TGTTAGGTGG CTGTGGGAAT TGCACTGCCA GCAGGCAGGA GGAATTCTGG GAGATGAAAT
1681    GGGATTGGGC AAGACCATCC AGATAATTGC CTTCTTGGCA GGTCTGAGCT ACAGCAAGAT
1741    CAGGACTCGT GGTTCAAATT ACAGGTTTGA GGGGTTGGGT CCAACTGTAA TTGTCTGTCC
1801    AACAACAGTG ATGCATCAGT GGGTGAAGGA ATTTCACACG TGGTGGCCTC CGTTCAGAGT
1861    GGCAATTCTA CATGAAACCG GTTCCTATAC CCACAAAAAG GAGAAACTAA TTCGAGATGT
1921    TGCTCATTGT CATGGAATTT TGATCACATC TTACTCCTAC ATTCGATTGA TGCAGGATGA
1981    CATTAGCAGG TATGACTGGC ACTATGTGAT CTTGGACGAA GGACACAAAA TTCGAAATCC
2041    AAATGCTGCT GTCACCCTTG CTTGCAAACA GTTTCGCACC CCTCATCGGA TCATTCTGTC
2101    TGGCTCACCG ATGCAAAATA ACCTCCGAGA GCTGTGGTCG CTCTTTGACT TCATCTTCCC
2161    GGGAAAGTTA GGCACGTTGC CTGTGTTTAT GGAGCAGTTC TCCGTCCCCA TCACCATGGG
2221    GGGATATTCA AATGCTTCCC CAGTACAGGT CAAAACTGCT TACAAGTGTG CATGTGTCTT
2281    ACGAGATACC ATAAATCCAT ACCTACTGCG GAGAATGAAG TCAGATGTCA AGATGAGCCT
2341    TTCTTTGCCA GATAAAAATG AACAGGTCTT ATTTTGCCGT CTTACAGATG AGCAGCATAA
2401    AGTCTACCAA AATTTCGTTG ATTCCAAAGA AGTTTACAGG ATTCTCAATG GAGAGATGCA
2461    GATTTTCTCC GGACTTATAG CCCTAAGAAA AATTTGCAAC CACCCTGATC TCTTTTCTGG
2521    AGGTCCCAAG AATCTCAAAG GTCTTCCTGA TGATGAACTA GAAGAAGATC AGTTTGGGTA
2581    CTGGAAACGT TCTGGGAAAA TGATTGTTGT TGAGTCTTTG TTGAAAATAT GGCACAAGCA
2641    GGGTCAGCGA GTATTGCTGT TTTCTCAGTC AAGGCAGATG CTGGACATAC TTGAAGTATT
2701    CCTTAGAGCC CAAAAGTATA CCTATCTCAA GATGGATGGT ACCACTACAA TAGCTTCAAG
2761    ACAGCCACTG ATTACGAGAT ACAATGAGGA CACATCCATA TTTGTGTTTC TTCTGACCAC
2821    GCGGGTGGGC GGCTTAGGTG TCAACCTGAC GGGGGCAAAC AGAGTTGTCA TCTATGACCC
2881    AGACTGGAAC CCAAGCACGG ACACGCAGGC CCGGGAGCGA GCATGGAGAA TAGGCCAGAA
2941    GAAGCAAGTG ACTGTGTACA GGCTCCTGAC TGCGGGCACC ATTGAAGAAA AGATCTACCA
3001    CCGACAAATC TTCAAGCAGT TTTTGACAAA TAGAGTGCTA AAAGACCCAA AACAAAGGCG
3061    GTTTTTCAAA TCCAATGATC TCTATGAGCT ATTTACTCTG ACTAGTCCTG ATGCATCCCA
3121    GAGCACTGAA ACAAGTGCAA TTTTTGCAGG AACTGGATCA GATGTTCAGA CACCCAAATG
3181    CCATCTAAAA AGAAGGATTC AACCAGCCTT TGGAGCAGAC CATGATGTTC CAAAACGCAA
3241    GAAGTTCCCT GCTTCTAACA TATCTGTAAA TGATGCCACA TCATCTGAAG AGAAATCTGA
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Please delete the disclosure on page 49 and replace it with the following:

DEFINITION Human mRNA for XPAC protein.(XPA) **(SEQ ID NO: 13)**
ACCESSION D14533

BASE COUNT 458 a 232 c 358 g 329 t
ORIGIN Chromosome 9.

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1       AGCTAGGTCC TCGGAGTGGG CCAGAGATGG CGGCGGCCGA CGGGGCTTTG CCGGAGGCGG
61      CGGCTTTAGA GCAACCCGCG GAGCTGCCTG CCTCGGTGCG GGCGAGTATC GAGCGGAAGC
121     GGCAGCGGGC ACTGATGCTG CGCCAGGCCC GGCTGGCTGC CCGGCCCTAC TCGGCGACGG
181     CGGCTGCGGC TACTGGAGGC ATGGCTAATG TAAAAGCAGC CCCAAAGATA ATTGACACAG
241     GAGGAGGCTT CATTTTAGAA GAGGAAGAAG AAGAAGAACA GAAAATTGGA AAAGTTGTTC
301     ATCAACCAGG ACCTGTTATG GAATTTGATT ATGTAATATG CGAAGAATGT GGGAAAGAAT
361     TTATGGATTC TTATCTTATG AACCACTTTG ATTTGCCAAC TTGTGATAAC TGCAGAGATG
421     CTGATGATAA ACACAAGCTT ATAACCAAAA CAGAGGCAAA ACAAGAATAT CTTCTGAAAG
481     ACTGTGATTT AGAAAAAAGA GAGCCACCTC TTAAATTTAT TGTGAAGAAG AATCCACATC
541     ATTCACAATG GGGTGATATG AAACTCTACT TAAAGTTACA GATTGTGAAG AGGTCTCTTG
601     AAGTTTGGGG TAGTCAAGAA GCATTAGAAG AAGCAAAGGA AGTCCGACAG GAAAACCGAG
661     AAAAAATGAA ACAGAAGAAA TTTGATAAAA AAGTAAAAGA ATTGCGGCGA GCAGTAAGAA
721     GCAGCGTGTG GAAAAGGGAG ACGATTGTTC ATCAACATGA GTATGGACCA GAAGAAAACC
781     TAGAAGATGA CATGTACCGT AAGACTTGTA CTATGTGTGG CCATGAACTG ACATATGAAA
841     AAATGTGATT TTTTAGTTCA GTGACCTGTT TTATAGAATT TTATATTTAA ATAAAGGAAA
901     TTTAGATTGG TCCTTTTCAA AATTCAAAAA AAAAAGCAAC ATCTTCATAG ATGAATGAAA
961     CCCTTGTATA AGTAATACTT CAGTAATAAT TATGTATGTT ATGGCTTAAA AGCAAGTTTC
1021    AGTGAAGGTC ACCTGGCCTG GTTGTGTGCA CAATGTCATG TCTGTGATTG CCTTCTTACA
1081    ACAGAGATGG GAGCTGAGTG CTAGAGTAGG TGCAGAAGTG GTAGGTCAGC TACAAATTTG
1141    AGGACAAGAT ACCAAGGCAA ACCCTAGATT GGGGTAGAGG GAAAAGGGTT CAACAAAGGC
1201    TGAACTGGAT TCTTAACCAA GAAACAAATA ATAGCAATGG TGGTGCACCA CTGTACCCCA
1261    GGTTCTAGTC ATGTGTTTTT TAGGACGATT TCTGTCTCCA CGATGGTGGA AACAGTGGGG
1321    AACTACTGCT GGAAAAAGCC CTAATAGCAG AAATAAACAT TGAGTTGTAC GAGTCTG
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